



and/or A

SEQUENCE LISTING

<111> Sheehan, John
Thomas, Daryl
Barrett, Wade

<120> Algorithms for selection of primer pairs

<130> 1011U1

<140> 10/042,406

<141> 2002-01-09

<150> 60/317,311

<151> 2001-09-05

<160> 2

<170> FastSEQ for Windows Version 4.0

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<212> DNA

<213> Human

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gacttaaata gataa 855
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<210> 2

<211> 855

<212> DNA

<213> Human

<220>

<221> misc_feature

<222> (306)...(433)

<223> n = A,T,C or G

<221> misc_feature

<222> (444)...(607)

<223> n = A,T,C or G

<400> 2

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gacttaaata gataa 855
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